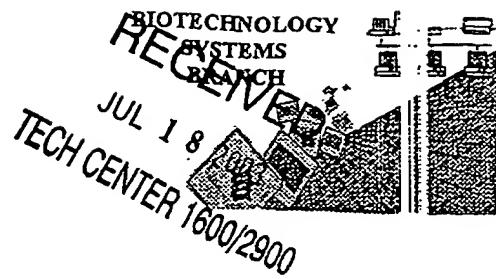


RAW SEQUENCE LISTING
ERROR REPORT



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/939,226 A
Source: 1600
Date Processed by STIC: 7/9/2003

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
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FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

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Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom. Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses

1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>>), EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry directly to:
U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202
Or
U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
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Revised 04/24/2003

03 SEP 13 5:30:10



1600

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/939,226A

DATE: 07/09/2003

TIME: 15:53:54

Input Set : A:\sequencelisting.txt

Output Set: N:\CRF4\07092003\I939226A.raw

3 <110> APPLICANT: Euroscreen S.A.
 4 SAMSON, Michel
 5 PARMENTIER, Marc
 6 VASSART, Gilbert
 7 LIBERT, Frederick
 9 <120> TITLE OF INVENTION: Methods for Identifying Compounds which Bind the Active CCR5
 10 Chemokine Receptor
 12 <130> FILE REFERENCE: 9409/2023C
 14 <140> CURRENT APPLICATION NUMBER: 09/939,226A
 15 <141> CURRENT FILING DATE: 2001-08-24
 17 <150> PRIOR APPLICATION NUMBER: US 08/833,752
 18 <151> PRIOR FILING DATE: 1997-04-09
 20 <150> PRIOR APPLICATION NUMBER: US 09/626,939
 21 <151> PRIOR FILING DATE: 2000-07-27
 23 <150> PRIOR APPLICATION NUMBER: US 08/810,028
 24 <151> PRIOR FILING DATE: 1997-03-04
 26 <160> NUMBER OF SEQ ID NOS: 18
 28 <170> SOFTWARE: PatentIn version 3.2
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 31 <211> LENGTH: 792
 32 <212> TYPE: DNA
 33 <213> ORGANISM: Homo sapiens
 35 <400> SEQUENCE: 1
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 38 tcccttact acaaaactc attgtttggc caaaaagaga gttaaattcaa tgttagacatc 120
 40 tatgttagca attaaaaacc tattgtatgt taaaaacagtt tgcattcatg gagggcaact 180
 42 aaatacattt taggacttta taaaatgtca ctttttattt atgcacaggg tggaaacaaga 240
 44 tggattatca agtgtcaatg ccaatctatg acatcaatc ttatacatcg gagccctgcc 300
 46 aaaaaatcaa tggaaagcaa atcgcagccc gctctctggcc tocgctctac tcactgggt 360
 48 tcatactttgg ttttggggc aacatgtctgg tcatactcat cctgataaaac tgcaaaaaggc 420
 50 tgaagagcat gactgacatc tacatgtctca acctggccat ctctgaccctg ttttccttc 480
 52 ttactgtccc cttctggct cactatgtc ccggccactg ggactttggaa aatacaatgt 540
 54 gtcaactctt gacagggctc tattttatag gcttcttctc tggaaatcttc ttcatcatcc 600
 56 tcctgacaat cgataggtac ctggctgtcg tccatgtgt gtttgccttta aaagccagga 660
 58 cggtcacctt tgggggtgggt acaagtgtga tcacttgggt ggtggctgtg tttgcgtctc 720
 60 tcccaggaat catcttacc agatctcaaa aagaaggctc tcattacacc tgcaagcttc 780
 62 attttccata ca 792
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 66 <211> LENGTH: 1477
 67 <212> TYPE: DNA
 68 <213> ORGANISM: Homo sapiens
 71 <220> FEATURE:
 72 <221> NAME/KEY: misc_feature

MP 69

Does Not Comply
Corrected Diskette Needed

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/939,226A

DATE: 07/09/2003
TIME: 15:53:54

Input Set : A:\sequencelistning.txt
Output Set: N:\CRF4\07092003\I939226A.raw

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74 <223> OTHER INFORMATION: Any nucleotide
76 <220> FEATURE:
77 <221> NAME/KEY: misc_feature
78 <222> LOCATION: (1384)..(1385)
79 <223> OTHER INFORMATION: Any nucleotide
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84 tcccttcaact acaaaaacttc attgcttgc caaaaagaga gtttaattcaa tgttagacatc 60
86 tatgttaggca attaaaaacc tattgtatgt taaaacagtt tgcatttcattg gaggcaact 120
88 aaatacattc taggacttta taaaagatca cttttattt atgcacaggg tggaaacaaga 180
90 tggattatca agtgtcaagt ccaatctatg acatcaatta ttatacatcg gagccctgcc 240
92 aaaaaatcaa tgtgaagcaa atcgcagccc gcctcctgcc tccgcctcac tcactggtgt 300
94 tcatactttgg ttttggc aacatgtgg tcatacctcat cctgataaaac tgcaaaaggc 360
96 tgaagagcat gactgacatc tacctgttca acctggccat ctctgacatc tttttccatc 420
98 ttactgtccc ttctggcact cactatgtgg cgcggcaggc ggactttggaaatacaatgt 480
100 gtcaacttttt gacaggcgtc tattttatag gtttttttc tggaaatcttc ttcatcatcc 540
102 tcctgacaat cgataggta ctggctgtcg tccatgtgt gtttgcattt aaaggcaggaa 600
104 cggtcaccc ttgggtgggt acaagtgtga tcacttgggt ggtggctgtg tttgcgtctc 660
106 tcccaggaaat catcttacc agatctcaaa aagaaggct tcattacacc tgcagctctc 720
108 attttccata cagtcgtat caattcttggaa agaatttcca gacattaaag atatgtcatct 780
110 tggggctgtt cctgcccgtg cttgtcatgg tcatacgttca ctgcggaaatc ctaaaaactc 840
112 tgcttgcgtt tcgaaatggaa aagaagaggc acagggtgtt gaggcttatac ttccatca 900
114 tgattgttta ttttcttcc tgggctccct acaacattgt cttttccctt aacaccccttcc 960
116 aggaattttt tggcctgaat aatttgcgtt gctctaacag gttggccaa gctatgcagg 1020
118 tgacagagac tcttggatg acgcactgtt gcatcaaccc catcatctat gccttgcgt 1080
120 gggagaagtt cagaaactac ctcttagtct ttttttttttcaaaatc gacattggcc aaacattttt 1140
122 gcaaatgttgc ttcttattttt cagcaagagg ctccggagcg agcaagctca gtttacaccc 1200
124 gatccactgg ggagcaggaa atatctgtgg gtttgcgttca gggactcaag tgggtgggt 1260
126 acccagtctt agtttgcac atggcttagt tttcatacac acgctgggtt ggggtnggt 1320
128 tggggatggt ttttttttttcaaaatc gacatttttca gtttgcgttca ggggtnggt 1380
130 atttggcatac ttttttttttcaaaatc gacatttttca gtttgcgttca ggggtnggt 1440
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134 <211> LENGTH: 1442
135 <212> TYPE: DNA
136 <213> ORGANISM: Homo sapiens
138 <400> SEQUENCE: 3
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141 tcccttcaact acaaaaacttc attgcttgc caaaaagaga gtttaattcaa tgttagacatc 60
143 tatgttaggca attaaaaacc tattgtatgt taaaacagtt tgcatttcattg gaggcaact 120
145 aaatacattc taggacttta taaaagatca cttttattt atgcacaggg tggaaacaaga 180
147 tggattatca agtgtcaagt ccaatctatg acatcaatta ttatacatcg gagccctgcc 240
149 aaaaaatcaa tgtgaagcaa atcgcagccc gcctcctgcc tccgcctcac tcactggtgt 300
151 tcatactttgg ttttggc aacatgtgg tcatacctcat cctgataaaac tgcaaaaggc 360
153 tgaagagcat gactgacatc tacctgttca acctggccat ctctgacatc tttttccatc 420
155 ttactgtccc ttctggcact cactatgtgg cgcggcaggc ggactttggaaatacaatgt 480
157 gtcaacttttt gacaggcgtc tattttatag gtttttttc tggaaatcc ttcatcatcc 540
159 tcctgacaat cgataggta ctggctgtcg tccatgtgtt gtttgcattt aaaggcaggaa 600
161 cggtcaccc ttgggtgggt acaagtgtga tcacttgggtt ggtggctgtg tttgcgtctc 660
162 tcatactttgg ttttggc aacatgtgg tcatacctcat cctgataaaac tgcaaaaggc 720

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/939,226A

DATE: 07/09/2003

TIME: 15:53:54

Input Set : A:\sequencelisting.txt

Output Set: N:\CRF4\07092003\I939226A.raw

163	tcccaggaat	catctttacc	agatctcaaa	aagaaggct	tcattacacc	tgca	gctctc	780									
165	atttccata	cattaaagat	agtcatctt	gggtctggcc	tgccgctgt	tgc	tcatggtc	840									
167	atctgtact	cggttacct	aaaaactct	cttcgggtgc	gaaatgagaa	gaagaggc	ac	900									
169	agggtgtga	ggcttatctt	caccatcatg	atttttatt	ttctcttctg	ggctccctac	960										
171	aacattgtcc	ttctcctgaa	caccccccag	gaatttctt	gcctgaataa	ttgc	actgtac	1020									
173	tctaaccagg	tggaccaagc	tatgcagg	acagagactc	ttggatgac	gcactgtgc	1080										
175	atcaacccca	tcatctatgc	cttgcggg	gagaagttca	gaaactact	cttagtctc	1140										
177	ttccaaaagc	acattgccaa	acgcttctgc	aaatgctgtt	ctatttcca	gcaagaggct	1200										
179	cccgagcgag	caagctca	ttacacccga	tccactgggg	agcaggaat	atctgtggc	1260										
181	ttgtgacacg	gactcaagt	ggctggtgac	ccagtcaag	ttgtgcacat	ggcttagtt	1320										
183	tcatacacag	cctgggctgg	gggtgggtgg	gaggctttt	ttaaaagaa	gttactgtta	1380										
185	tagagggct	aagattcata	catttattt	gcatctgtt	aaagtagatt	agatccgaat	1440										
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191	<211>	LENGTH:	184														
192	<212>	TYPE:	PRT														
193	<213>	ORGANISM:	Homo sapiens														
195	<400>	SEQUENCE:	4														
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198	1			5					10				15				
201	Ser	Glu	Pro	Cys	Gln	Lys	Ile	Asn	Val	Lys	Gln	Ile	Ala	Ala	Arg	Leu	
202				20					25				30				
205	Leu	Pro	Pro	Leu	Tyr	Ser	Leu	Val	Phe	Ile	Phe	Gly	Phe	Val	Gly	Asn	
206				35					40				45				
209	Met	Leu	Val	Ile	Leu	Ile	Leu	Ile	Asn	Cys	Lys	Arg	Leu	Lys	Ser	Met	
210				50					55				60				
213	Thr	Asp	Ile	Tyr	Leu	Leu	Asn	Leu	Ala	Ile	Ser	Asp	Leu	Phe	Phe	Leu	
214				65					70				75				80
217	Leu	Thr	Val	Pro	Phe	Trp	Ala	His	Tyr	Ala	Ala	Ala	Gln	Trp	Asp	Phe	
218				85					90				95				
221	Gly	Asn	Thr	Met	Cys	Gln	Leu	Leu	Thr	Gly	Leu	Tyr	Phe	Ile	Gly	Phe	
222				100					105				110				
225	Phe	Ser	Gly	Ile	Phe	Phe	Ile	Ile	Leu	Leu	Thr	Ile	Asp	Arg	Tyr	Leu	
226				115					120				125				
229	Ala	Val	Val	His	Ala	Val	Phe	Ala	Leu	Lys	Ala	Arg	Thr	Val	Thr	Phe	
230				130					135				140				
233	Gly	Val	Val	Thr	Ser	Val	Ile	Thr	Trp	Val	Val	Ala	Val	Phe	Ala	Ser	
234				145					150				155				160
237	Leu	Pro	Gly	Ile	Ile	Phe	Thr	Arg	Ser	Gln	Lys	Glu	Gly	Leu	His	Tyr	
238				165					170				175				
241	Thr	Cys	Ser	Ser	His	Phe	Pro	Tyr									
242				180													
245	<210>	SEQ ID NO:	5														
246	<211>	LENGTH:	352														
247	<212>	TYPE:	PRT														
248	<213>	ORGANISM:	Homo sapiens														
250	<400>	SEQUENCE:	5														
252	Met	Asp	Tyr	Gln	Val	Ser	Ser	Pro	Ile	Tyr	Asp	Ile	Asn	Tyr	Tyr	Thr	
253	1			5					10				15				

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/939,226A

DATE: 07/09/2003
TIME: 15:53:54

Input Set : A:\sequencelisting.txt
Output Set: N:\CRF4\07092003\I939226A.raw

256 Ser Glu Pro Cys Gln Lys Ile Asn Val Lys Gln Ile Ala Ala Arg Leu
257 20 25 30
260 Leu Pro Pro Leu Tyr Ser Leu Val Phe Ile Phe Gly Phe Val Gly Asn
261 35 40 45
264 Met Leu Val Ile Leu Ile Leu Ile Asn Cys Lys Arg Leu Lys Ser Met
265 50 55 60
268 Thr Asp Ile Tyr Leu Leu Asn Leu Ala Ile Ser Asp Leu Phe Phe Leu
269 65 70 75 80
272 Leu Thr Val Pro Phe Trp Ala His Tyr Ala Ala Ala Gln Trp Asp Phe
273 85 90 95
276 Gly Asn Thr Met Cys Gln Leu Leu Thr Gly Leu Tyr Phe Ile Gly Phe
277 100 105 110
280 Phe Ser Gly Ile Phe Phe Ile Leu Leu Thr Ile Asp Arg Tyr Leu
281 115 120 125
284 Ala Val Val His Ala Val Phe Ala Leu Lys Ala Arg Thr Val Thr Phe
285 130 135 140
288 Gly Val Val Thr Ser Val Ile Thr Trp Val Val Ala Val Phe Ala Ser
289 145 150 155 160
292 Leu Pro Gly Ile Ile Phe Thr Arg Ser Gln Lys Glu Gly Leu His Tyr
293 165 170 175
296 Thr Cys Ser Ser His Phe Pro Tyr Ser Gln Tyr Gln Phe Trp Lys Asn
297 180 185 190
300 Phe Gln Thr Leu Lys Ile Val Ile Leu Gly Leu Val Leu Pro Leu Leu
301 195 200 205
304 Val Met Val Ile Cys Tyr Ser Gly Ile Leu Lys Thr Leu Leu Arg Cys
305 210 215 220
308 Arg Asn Glu Lys Lys Arg His Arg Ala Val Arg Leu Ile Phe Thr Ile
309 225 230 235 240
312 Met Ile Val Tyr Phe Leu Phe Trp Ala Pro Tyr Asn Ile Val Leu Leu
313 245 250 255
316 Leu Asn Thr Phe Gln Glu Phe Phe Gly Leu Asn Asn Cys Ser Ser Ser
317 260 265 270
320 Asn Arg Leu Asp Gln Ala Met Gln Val Thr Glu Thr Leu Gly Met Thr
321 275 280 285
324 His Cys Cys Ile Asn Pro Ile Ile Tyr Ala Phe Val Gln Glu Lys Phe
325 290 295 300
328 Arg Asn Tyr Leu Leu Val Phe Phe Gln Lys His Ile Ala Lys Arg Phe
329 305 310 315 320
332 Cys Lys Cys Cys Ser Ile Phe Gln Gln Glu Ala Pro Glu Arg Ala Ser
333 325 330 335
336 Ser Val Tyr Thr Arg Ser Thr Gly Glu Gln Glu Ile Ser Val Gly Leu
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340 <210> SEQ ID NO: 6
341 <211> LENGTH: 215
342 <212> TYPE: PRT
343 <213> ORGANISM: Homo sapiens
345 <400> SEQUENCE: 6
347 Met Asp Tyr Gln Val Ser Ser Pro Ile Tyr Asp Ile Asn Tyr Tyr Thr
348 1 5 10 15

RAW SEQUENCE LISTING

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TIME: 15:53:54

Input Set : A:\sequencelisting.txt

Output Set: N:\CRF4\07092003\I939226A.raw

351 Ser Glu Pro Cys Gln Lys Ile Asn Val Lys Gln Ile Ala Ala Arg Leu
 352 20 25 30
 355 Leu Pro Pro Leu Tyr Ser Leu Val Phe Ile Phe Gly Phe Val Gly Asn
 356 35 40 45
 359 Met Leu Val Ile Leu Ile Leu Ile Asn Cys Lys Arg Leu Lys Ser Met
 360 50 55 60
 363 Thr Asp Ile Tyr Leu Leu Asn Leu Ala Ile Ser Asp Leu Phe Phe Leu
 364 65 70 75 80
 367 Leu Thr Val Pro Phe Trp Ala His Tyr Ala Ala Ala Gln Trp Asp Phe
 368 85 90 95
 371 Gly Asn Thr Met Cys Gln Leu Leu Thr Gly Leu Tyr Phe Ile Gly Phe
 372 100 105 110
 375 Phe Ser Gly Ile Phe Phe Ile Ile Leu Leu Thr Ile Asp Arg Tyr Leu
 376 115 120 125
 379 Ala Val Val His Ala Val Phe Ala Leu Lys Ala Arg Thr Val Thr Phe
 380 130 135 140
 383 Gly Val Val Thr Ser Val Ile Thr Trp Val Val Ala Val Phe Ala Ser
 384 145 150 155 160
 387 Leu Pro Gly Ile Ile Phe Thr Arg Ser Gln Lys Glu Gly Leu His Tyr
 388 165 170 175
 391 Thr Cys Ser Ser His Phe Pro Tyr Ile Lys Asp Ser His Leu Gly Ala
 392 180 185 190
 395 Gly Pro Ala Ala Ala Cys His Gly His Leu Leu Leu Gly Asn Pro Lys
 396 195 200 205
 399 Asn Ser Ala Ser Val Ser Lys
 400 210 215
 403 <210> SEQ ID NO: 7
 404 <211> LENGTH: 360
 405 <212> TYPE: PRT
 406 <213> ORGANISM: Homo sapiens
 409 <220> FEATURE:
 410 <221> NAME/KEY: MISC_FEATURE
 411 <222> LOCATION: (325)..(327)
 412 <223> OTHER INFORMATION: Xaa = any amino acid
 414 <400> SEQUENCE: 7
 416 Met Leu Ser Thr Ser Arg Ser Arg Phe Ile Arg Asn Thr Asn Glu Ser
 417 1 5 10 15
 420 Gly Glu Glu Val Thr Thr Phe Phe Asp Tyr Asp Tyr Gly Ala Pro Cys
 421 20 25 30
 424 His Lys Phe Asp Val Lys Gln Ile Gly Ala Gln Leu Leu Pro Pro Leu
 425 35 40 45
 428 Tyr Ser Leu Val Phe Ile Phe Gly Phe Val Gly Asn Met Leu Val Val
 429 50 55 60
 432 Leu Ile Leu Ile Asn Cys Lys Lys Leu Lys Cys Leu Thr Asp Ile Tyr
 433 65 70 75 80
 436 Leu Leu Asn Leu Ala Ile Ser Asp Leu Leu Phe Ile Ile Thr Leu Pro
 437 85 90 95
 440 Leu Trp Ala His Ser Ala Ala Asn Glu Trp Val Phe Gly Asn Ala Met
 441 100 105 110

09/939,226A 15

<210> SEQ ID NO 14
<211> LENGTH: 27
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: :
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see p.7 for error explanation.

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<210> SEQ ID NO 15
<211> LENGTH: 27
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
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<213> ORGANISM: Artificial Sequence
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28

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<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: :
<400> SEQUENCE: 17
ctgatctaga gccatgtgca caactct

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RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/09/939,226A

DATE: 07/09/2003
TIME: 15:53:55

Input Set : A:\sequencelisting.txt
Output Set: N:\CRF4\07092003\I939226A.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <210> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:2; N Pos. 1377,1384,1385
Seq#:7; Xaa Pos. 325,326,327
Seq#:8; Xaa Pos. 231,232,233,333,334,335
Seq#:10; Xaa Pos. 145,146,147,321,322,323

✓OK

Use of <220> Feature(NEW RULES):

Sequence(s) are missing the <220> Feature and associated headings.
Use of <220> to <223> is MANDATORY if <213> ORGANISM is "Artificial Sequence" or "Unknown". Please explain source of genetic material in <220> to <223> section (See "Federal Register;" 6/01/98, Vol. 63, No. 104, pp.29631-32) (Sec.1.823 of new Rules)

Seq#:14,15,16,17

T
error explanation

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/939,226A

DATE: 07/09/2003

TIME: 15:53:55

Input Set : A:\sequencelistning.txt

Output Set: N:\CRF4\07092003\I939226A.raw

L:126 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:1320
M 341 Repeated in SeqNo=2
L 496 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:320
L 582 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:224
M 341 Repeated in SeqNo=8
L:771 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:144
M:341 Repeated in SeqNo=10
L:885 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:14
L:887 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ#:14, <213>
ORGANISM:Artificial Sequence
L:887 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:14, <213>
ORGANISM:Artificial Sequence
L:887 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:14,Line#:887
L:894 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:15
L:896 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ#:15, <213>
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L:896 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:15, <213>
ORGANISM:Artificial Sequence
L:896 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:15,Line#:896
L:903 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:16
L:905 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ#:16, <213>
ORGANISM:Artificial Sequence
L:905 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:16, <213>
ORGANISM:Artificial Sequence
L:905 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:16,Line#:905
L:912 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:17
L:914 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ#:17, <213>
ORGANISM:Artificial Sequence
L:914 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:17, <213>
ORGANISM:Artificial Sequence
L:914 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:17,Line#:914